AMENDMENTS TO THE SPECIFICAITON

Page 6, lines 10-13, please amend as follows:

The amino acid sequence of the extracellular domain of human IL-10R was used to screen the database of the Sanger Center (http://www.sanger.ac.uk/e/cgi-bin/nph-Blast_server; html), incorporated by reference, using TBLASTN software.

Page 6, lines 18-21: please amend as follows:

Once the BAC sequence was identified, it was analyzed further, using the NIX analysis program found at http://www.hgmp.mrc.ac.uk/Registerd/Webapp/nix/, incorporated by reference. The software predicted a gene comprising 5 exons, stretching over about 16 kilobases, with the last exon corresponding to several EST sequences.